SEQUENCE LISTING

<110> Bacher, Adelbert
Mortl, Simone

<120> METHOD FOR SCREENING FOR INHIBITORS OF THE BIOSYNTHESIS OF RIBOFLAVIN

<130> 9286-9

<140> US 10/070,144
<141> 2002-03-01

<150> DE 199 42 175.7
<151> 1999-09-03

<160> 12

<170> PatentIn version 3.1

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<213> Artificial sequence

<220>
<223> Synthetic oligonucleotide primer

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Met Lys Ser Leu Ala Ser Pro Pro Cys Leu Arg Leu Ile Pro Thr Ala
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cac cgt cag ctc aat tcg cgt caa tct tcc tcc gcc tgt tat ata cac 96
His Arg Gln Leu Asn Ser Arg Gln Ser Ser Ser Ala Cys Tyr Ile His
20 25 30

ggt ggc tct tct gtg aac aaa tcc aat aat ctc tca ttc tcc tca tcc 144
Gly Gly Ser Ser Val Asn Lys Ser Asn Asn Leu Ser Phe Ser Ser Ser
35 40 45

aca tcc gga ttt gcg tca cca cta gct gta gag aag gaa tta cgc tct 192
Thr Ser Gly Phe Ala Ser Pro Leu Ala Val Glu Lys Glu Leu Arg Ser
50 55 60

tca ttc gta cag acg gct gct gtt cgc cat gtt acg ggg tct ctt atc 240
Ser Phe Val Gln Thr Ala Ala Val Arg His Val Thr Gly Ser Leu Ile
65 70 75 80

aga ggc gaa ggt ctt aga ttc gcc atc gtg gta gct cgt ttc aat gag 288
Arg Gly Glu Gly Leu Arg Phe Ala Ile Val Val Ala Arg Phe Asn Glu
85 90 95

gtt gtg act aag ttg ctt ttg gaa gga gcg att gag act ttc aag aag 336
Val Val Thr Lys Leu Leu Leu Glu Gly Ala Ile Glu Thr Phe Lys Lys
100 105 110

tat tca gtc aga gaa gaa gac att gaa gtt att tgg gtt cct ggc agc 384

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Tyr Ser Val Arg Glu Glu Asp Ile Glu Val Ile Trp Val Pro Gly Ser
        115                 120                 125

ttt gaa att ggt gtt gtt gca caa aat ctt ggg aaa tcg gga aaa ttt      432
Phe Glu Ile Gly Val Val Ala Gln Asn Leu Gly Lys Ser Gly Lys Phe
    130                 135                 140

cat gct gtt tta tgt atc ggc gct gtg ata aga gga gat acc aca cat      480
His Ala Val Leu Cys Ile Gly Ala Val Ile Arg Gly Asp Thr Thr His
145                 150                 155                 160

tat gat gct gtt gcc aac tct gct gcg tct gga gta ctt tct gct agc      528
Tyr Asp Ala Val Ala Asn Ser Ala Ala Ser Gly Val Leu Ser Ala Ser
                165                 170                 175

ata aat tca ggc gtt cca tgc ata ttt ggt gta ctg act tgc gag gac      576
Ile Asn Ser Gly Val Pro Cys Ile Phe Gly Val Leu Thr Cys Glu Asp
            180                 185                 190

atg gat cag gct ctg aat cga tct ggt ggc aaa gcc ggc aat aag gga      624
Met Asp Gln Ala Leu Asn Arg Ser Gly Gly Lys Ala Gly Asn Lys Gly
        195                 200                 205

gct gaa act gct ttg acg gcg ctc gaa atg gcg tcg ttg ttt gag cac      672
Ala Glu Thr Ala Leu Thr Ala Leu Glu Met Ala Ser Leu Phe Glu His
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His Arg Gln Leu Asn Ser Arg Gln Ser Ser Ser Ala Cys Tyr Ile His
            20                  25                  30

Gly Gly Ser Ser Val Asn Lys Ser Asn Asn Leu Ser Phe Ser Ser Ser
        35                  40                  45

Thr Ser Gly Phe Ala Ser Pro Leu Ala Val Glu Lys Glu Leu Arg Ser
    50                  55                  60

Ser Phe Val Gln Thr Ala Ala Val Arg His Val Thr Gly Ser Leu Ile
65                  70                  75                  80

Arg Gly Glu Gly Leu Arg Phe Ala Ile Val Val Ala Arg Phe Asn Glu
                85                  90                  95
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Val Val Thr Lys Leu Leu Leu Glu Gly Ala Ile Glu Thr Phe Lys Lys
100 105 110

Tyr Ser Val Arg Glu Glu Asp Ile Glu Val Ile Trp Val Pro Gly Ser
115 120 125

Phe Glu Ile Gly Val Val Ala Gln Asn Leu Gly Lys Ser Gly Lys Phe
130 135 140

His Ala Val Leu Cys Ile Gly Ala Val Ile Arg Gly Asp Thr Thr His
145 150 155 160

Tyr Asp Ala Val Ala Asn Ser Ala Ala Ser Gly Val Leu Ser Ala Ser
165 170 175

Ile Asn Ser Gly Val Pro Cys Ile Phe Gly Val Leu Thr Cys Glu Asp
180 185 190

Met Asp Gln Ala Leu Asn Arg Ser Gly Gly Lys Ala Gly Asn Lys Gly
195 200 205

Ala Glu Thr Ala Leu Thr Ala Leu Glu Met Ala Ser Leu Phe Glu His
210 215 220

His Leu Lys
225

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Met Lys Ser Leu Ala Ser Pro Pro Cys Leu Arg Leu Ile Pro Thr Ala
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His Arg Gln Leu Asn Ser Arg Gln Ser Ser Ser Ala Cys Tyr Ile His
20 25 30

Gly Gly Ser Ser Val Asn Lys Ser Asn Asn Leu Ser Phe Ser Ser Ser
35 40 45

Thr Ser Gly Phe Ala Ser Pro Leu Ala Val Glu Lys Glu Leu Arg Ser
50 55 60

Ser Phe Val Gln Thr Ala Ala Val Arg His Val Thr Gly Ser Leu Ile
65 70 75 80

Arg Gly Glu Gly Leu Arg Phe Ala Ile Val Val Ala Arg Phe Asn Glu
85 90 95

Val Val Thr Lys Leu Leu Leu Glu Gly Ala Ile Glu Thr Phe Lys Lys
100 105 110

Tyr Ser

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Val Arg Glu Glu Asp Ile Glu Val Ile Trp Val Pro Gly Ser Phe Glu
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Ile Gly Val Val Ala Gln Asn Leu Gly Lys Ser Gly Lys Phe His Ala
20 25 30

Val Leu Cys Ile Gly Ala Val Ile Arg Gly Asp Thr Thr His Tyr Asp
35 40 45

Ala Val Ala Asn Ser Ala Ala Ser Gly Val Leu Ser Ala Ser Ile Asn
50 55 60

Ser Gly Val Pro Cys Ile Phe Gly Val Leu Thr Cys Glu Asp Met Asp
65 70 75 80

Gln Gln Ala Leu Asn Arg Ser Gly Gly Lys Ala Gly Asn Lys Gly Ala
85 90 95

Glu Thr Ala Leu Thr Ala Leu Glu Met Ala Ser Leu Phe Glu His His
100 105 110

Leu Lys

<210> 10
<211> 156
<212> PRT
<213> Escherichia coli

<400> 10

Met Asn Ile Ile Glu Ala Asn Val Ala Thr Pro Asp Ala Arg Val Ala
1 5 10 15

Ile Thr Ile Ala Arg Phe Asn Asn Phe Ile Asn Asp Ser Leu Leu Glu
20 25 30

Gly Ala Ile Asp Ala Leu Lys Arg Ile Gly Gln Val Lys Asp Glu Asn
35 40 45

Ile Thr Val Val Trp Val Pro Gly Ala Tyr Glu Leu Pro Leu Ala Ala
50 55 60

Gly Ala Leu Ala Lys Thr Gly Lys Tyr Asp Ala Val Ile Ala Leu Gly
65 70 75 80

Thr Val Ile Arg Gly Gly Thr Ala His Phe Glu Tyr Val Ala Gly Gly
85 90 95

Ala Ser Asn Gly Leu Ala His Val Ala Gln Asp Ser Glu Ile Pro Val
100 105 110

Ala Phe Gly Val Leu Thr Thr Glu Ser Ile Glu Gln Ala Ile Glu Arg
115 120 125

Ala Gly Thr Lys Ala Gly Asn Lys Gly Ala Glu Ala Ala Leu Thr Ala
130 135 140

Leu Glu Met Ile Asn Val Leu Lys Ala Ile Lys Ala
145 150 155

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<213> Bacillus subtilis

<400> 11

Met Asn Ile Ile Gln Gly Asn Leu Val Gly Thr Gly Leu Lys Ile Gly
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Ile Val Val Gly Arg Phe Asn Asp Phe Ile Thr Ser Lys Leu Leu Ser
20 25 30

Gly Ala Glu Asp Ala Leu Leu Arg His Gly
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<213> Bacillus subtilis

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Ile Pro Phe Ala Ala Lys Lys Met Ala Glu Thr Lys Lys Tyr Asp Ala
            20                  25                  30

Ile Ile Thr Leu Gly Thr Val Ile Arg Gly Ala Thr Thr His Tyr Asp
        35                  40                  45

Tyr Val Cys Asn Glu Ala Ala Lys Gly Ile Ala Gln Ala Ala Asn Thr
    50                  55                  60

Thr Gly Val Pro Val Ile Phe Gly Ile Val Thr Thr Glu Asn Ile Glu
65                  70                  75                  80

Gln Ala Ile Glu Arg Ala Gly Thr Lys Ala Gly Asn Lys Gly Val Asp
                85                  90                  95

Cys Ala Val Ser Ala Ile Glu Met Ala Asn Leu Asn Arg Ser Phe Glu
            100                 105                 110
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